

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/019,282
Source: PCT/10
Date Processed by STIC: 10/19/04

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 10/19/2004

PATENT APPLICATION: US/10/019,282

TIME: 17:29:38

Input Set : D:\217770US0PCT.txt

Output Set: N:\CRF4\10192004\J019282.raw

THE

```

3 <110> APPLICANT: SHIBATA, TAKASHI
4     NOGUCHI, YUJI
5     YAMASHITA, MICHIO
7 <120> TITLE OF INVENTION: GENE ENCODING CYCLIC LIPOPEPTIDE ACYLASE AND EXPRESSION OF
8     SAME
10 <130> FILE REFERENCE: 217770US0PCT
12 <140> CURRENT APPLICATION NUMBER: 10/019,282
13 <141> CURRENT FILING DATE: 2002-01-02
15 <150> PRIOR APPLICATION NUMBER: PCT/JP00/04285
16 <151> PRIOR FILING DATE: 2000-06-28
18 <150> PRIOR APPLICATION NUMBER: JP189644/1999
19 <151> PRIOR FILING DATE: 1999-07-02
21 <160> NUMBER OF SEQ ID NOS: 76
23 <170> SOFTWARE: PatentIn version 3.2
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 5692
27 <212> TYPE: DNA
28 <213> ORGANISM: Streptomyces Sp.
31 <220> FEATURE:
32 <221> NAME/KEY: CDS
33 <222> LOCATION: (948)..(3362)
35 <400> SEQUENCE: 1
36 gaattccgga tggttggaga ggccgatcca gacggtgggc ggggcgaaga ggctgtcggc      60
38 caggcccgtc tgcagcaggt cgaagatcga ggcggcgctc ggaccgtcca ggatgggtgtt      120
40 ctccgcgcgc accgcagcat agggcagcag gaacacgtgc atctggggccg agtggtagag      180
42 cggcagggag tgcacggggc ggtcggctgc ggcgaggccg agcgcggtga tcgcgctgac      240
44 gtactcgtgg accagggccc cgtgcgtcat catcgcgccc ttgggcaggg cgggtggtccc      300
46 ggaggtgtac agcagctgca ccaggtcgtc ggaggcgggc gggcgccgcg ggggtgaacgc      360
48 ccgttccgtc tccagggcgt cgagcagcga gccgggcgcg tcgcggagcg cgcgcaccgg      420
50 ggtccgcgcg gggagccgcc cggcgaggtc cgggtcggtc aggacgaggg aggagccgga      480
52 ctggtcagag aggtaggcca ggtcgtcgcc ggtgaggttc tggttgaccg gtacgtggac      540
54 gagaccggcc cgtgcgcagg cgaggaagcc gatcagatag gcgtcggagt tgtgcgcgta      600
56 ggcggccacc cggtcgcgcg gggcgagagc gtactcctcg gtgaggacgg cggcggccgt      660
58 ggagacggcg gcgtccaggg agcggtaggt ccaggtccgg tcggcgtagc gcacggcggt      720
60 ccggtcgggg gtgcgcgggg cgctgcgggt gaggacgccg tcgactgtgc tgctgcgtac      780
62 acctgtcatg gcgtgatcct gtgcgtccgg gccctcgggg gtcaagaggc tggataaccga      840
64 ccagacgggt gacagcttcc cgggctccct ggctgagtga cgcttgggccg tccgggcggt      900
66 ccggaaccgg cgcgcccgtg ccaccctgac cgctgggagg aaacacc ttg acg tta      956
67                                     Leu Thr Leu
68                                     1
70 cgc aac cgt ctg aga ctg ctc ggg gtc gcc ggt ctc gcc ctg ttc acc      1004
71 Arg Asn Arg Leu Arg Leu Leu Gly Val Ala Gly Leu Ala Leu Phe Thr
72      5              10              15

```

RAW SEQUENCE LISTING

DATE: 10/19/2004

PATENT APPLICATION: US/10/019,282

TIME: 17:29:38

Input Set : D:\217770US0PCT.txt

Output Set: N:\CRF4\10192004\J019282.raw

74	gtg	tcg	gcg	tcg	ctg	ccg	cct	gcc	acc	gcg	tcc	ggg	acc	cag	gag	acg	1052
75	Val	Ser	Ala	Ser	Leu	Pro	Pro	Ala	Thr	Ala	Ser	Gly	Thr	Gln	Glu	Thr	
76	20					25				30					35		
78	cgg	cac	ccg	tcc	ggg	agc	ggg	ctt	tcg	gcc	gtc	atc	cgg	tac	acg	gag	1100
79	Arg	His	Pro	Ser	Gly	Ser	Gly	Leu	Ser	Ala	Val	Ile	Arg	Tyr	Thr	Glu	
80				40				45						50			
82	tac	ggc	att	ccg	cac	atc	gtg	gcg	gag	gac	tac	gcg	cag	ttg	ggc	ttc	1148
83	Tyr	Gly	Ile	Pro	His	Ile	Val	Ala	Glu	Asp	Tyr	Ala	Gln	Leu	Gly	Phe	
84			55					60					65				
86	ggc	acc	ggc	tgg	gcg	cag	gcc	gcc	gat	cag	gtg	tgc	acg	ctg	gcg	gac	1196
87	Gly	Thr	Gly	Trp	Ala	Gln	Ala	Ala	Asp	Gln	Val	Cys	Thr	Leu	Ala	Asp	
88			70					75					80				
90	ggc	ttc	ctc	acg	gtg	cgc	ggg	gag	cgg	tcg	agg	ttc	ttc	ggc	ccg	gac	1244
91	Gly	Phe	Leu	Thr	Val	Arg	Gly	Glu	Arg	Ser	Arg	Phe	Phe	Gly	Pro	Asp	
92		85				90						95					
94	gcc	gcc	acg	gac	tac	tcc	ctc	tcc	tcg	gcg	gcg	acg	aac	ctc	tcc	agc	1292
95	Ala	Ala	Thr	Asp	Tyr	Ser	Leu	Ser	Ser	Ala	Ala	Thr	Asn	Leu	Ser	Ser	
96	100					105					110					115	
98	gac	ctg	tac	ttc	cgg	ggc	gtc	cgc	gac	agc	ggc	acg	gtg	gag	aag	ctg	1340
99	Asp	Leu	Tyr	Phe	Arg	Gly	Val	Arg	Asp	Ser	Gly	Thr	Val	Glu	Lys	Leu	
100				120						125					130		
102	ctc	aag	gag	ccc	gcg	ccc	gcc	ggg	ccg	agc	agg	gac	gtc	aag	gag	acg	1388
103	Leu	Lys	Glu	Pro	Ala	Pro	Ala	Gly	Pro	Ser	Arg	Asp	Val	Lys	Glu	Thr	
104				135					140					145			
106	atg	cgc	ggg	ttc	gcc	gcc	ggg	tac	aac	gcg	tgg	atc	gcg	cag	aac	cgg	1436
107	Met	Arg	Gly	Phe	Ala	Ala	Gly	Tyr	Asn	Ala	Trp	Ile	Ala	Gln	Asn	Arg	
108			150					155					160				
110	atc	acc	gac	ccc	gcc	tgc	cgg	ggc	gcg	tcc	tgg	gtg	cgc	ccg	gtg	acg	1484
111	Ile	Thr	Asp	Pro	Ala	Cys	Arg	Gly	Ala	Ser	Trp	Val	Arg	Pro	Val	Thr	
112		165					170						175				
114	gcg	ctg	gac	gtg	gcg	gcg	cgc	ggc	tac	gcg	ctg	gcg	gtg	ctc	ggc	ggc	1532
115	Ala	Leu	Asp	Val	Ala	Ala	Arg	Gly	Tyr	Ala	Leu	Ala	Val	Leu	Gly	Gly	
116	180					185					190				195		
118	cag	ggg	cgc	ggc	atc	gac	ggc	atc	acc	gcg	gca	cag	ccg	ccg	acc	gcc	1580
119	Gln	Gly	Arg	Gly	Ile	Asp	Gly	Ile	Thr	Ala	Ala	Gln	Pro	Pro	Thr	Ala	
120				200						205					210		
122	gct	cct	ccg	gcg	gcc	ggg	gtc	acg	ccc	gag	gag	gcg	gcg	acg	gcg	gcg	1628
123	Ala	Pro	Pro	Ala	Ala	Gly	Val	Thr	Pro	Glu	Glu	Ala	Ala	Thr	Ala	Ala	
124				215						220				225			
126	gag	cgg	ctg	ctg	tcg	acg	cag	aac	gcg	gac	atg	ggg	tcc	aac	gcg	gtg	1676
127	Glu	Arg	Leu	Leu	Ser	Thr	Gln	Asn	Ala	Asp	Met	Gly	Ser	Asn	Ala	Val	
128			230					235					240				
130	gcc	ttc	gac	ggc	tcc	acg	acg	gtg	aac	ggg	cgc	ggg	ctg	ttg	ctc	ggc	1724
131	Ala	Phe	Asp	Gly	Ser	Thr	Thr	Val	Asn	Gly	Arg	Gly	Leu	Leu	Leu	Gly	
132		245					250					255					
134	aac	ccg	cac	tac	ccg	tgg	cag	ggc	gga	cgc	cgc	ttc	tgg	cag	gcg	cag	1772
135	Asn	Pro	His	Tyr	Pro	Trp	Gln	Gly	Gly	Arg	Arg	Phe	Trp	Gln	Ala	Gln	
136	260					265					270					275	
138	cag	acg	atc	ccc	ggc	gag	ctg	aac	gtg	tcg	ggc	gcg	tcc	ctg	ctg	ggc	1820

RAW SEQUENCE LISTING

DATE: 10/19/2004

PATENT APPLICATION: US/10/019,282

TIME: 17:29:38

Input Set : D:\217770USOPCT.txt

Output Set: N:\CRF4\10192004\J019282.raw

139	Gln	Thr	Ile	Pro	Gly	Glu	Leu	Asn	Val	Ser	Gly	Ala	Ser	Leu	Leu	Gly	
140					280					285					290		
142	gcg	acg	acg	atc	tcg	atc	ggg	cac	aac	gcc	gat	gtg	gcg	tgg	agc	cat	1868
143	Ala	Thr	Thr	Ile	Ser	Ile	Gly	His	Asn	Ala	Asp	Val	Ala	Trp	Ser	His	
144				295					300					305			
146	acg	gtc	gcc	acg	ggc	gtc	acg	ctg	aat	ctg	cat	cag	ctc	agc	ctc	gat	1916
147	Thr	Val	Ala	Thr	Gly	Val	Thr	Leu	Asn	Leu	His	Gln	Leu	Ser	Leu	Asp	
148			310					315					320				
150	ccg	gcc	gac	ccg	acc	gtc	tat	ctg	gtg	gac	ggg	aag	cgg	gag	cgg	atg	1964
151	Pro	Ala	Asp	Pro	Thr	Val	Tyr	Leu	Val	Asp	Gly	Lys	Arg	Glu	Arg	Met	
152		325					330					335					
154	acg	cag	cgg	acg	gtg	agc	gtc	ccg	gtg	aag	ggc	ggg	gcc	gac	gtg	acc	2012
155	Thr	Gln	Arg	Thr	Val	Ser	Val	Pro	Val	Lys	Gly	Gly	Ala	Asp	Val	Thr	
156	340					345					350					355	
158	cgc	acc	cag	tgg	tgg	acc	cgc	tac	ggg	ccg	gtg	gcc	acc	tcg	atg	ggc	2060
159	Arg	Thr	Gln	Trp	Trp	Thr	Arg	Tyr	Gly	Pro	Val	Ala	Thr	Ser	Met	Gly	
160				360					365					370			
162	gcg	ggg	ctg	ccg	ttg	ccg	tgg	acg	gcg	agc	acg	gcg	tac	gcg	ctg	aac	2108
163	Ala	Gly	Leu	Pro	Leu	Pro	Trp	Thr	Ala	Ser	Thr	Ala	Tyr	Ala	Leu	Asn	
164			375					380					385				
166	gat	ccg	aac	gcg	acg	aat	ctg	cgg	atg	gcg	gac	acc	ggt	ctg	ggc	ttc	2156
167	Asp	Pro	Asn	Ala	Thr	Asn	Leu	Arg	Met	Ala	Asp	Thr	Gly	Leu	Gly	Phe	
168		390					395					400					
170	ggc	aag	gcc	cgc	tcc	acg	ggt	gac	gtc	gag	cgt	gcg	ctg	cac	cgg	tcg	2204
171	Gly	Lys	Ala	Arg	Ser	Thr	Gly	Asp	Val	Glu	Arg	Ala	Leu	His	Arg	Ser	
172		405					410					415					
174	cag	ggc	atg	ccg	tgg	gtg	aac	acg	atc	gcg	gcg	gac	cgg	gcg	ggt	cgc	2252
175	Gln	Gly	Met	Pro	Trp	Val	Asn	Thr	Ile	Ala	Ala	Asp	Arg	Ala	Gly	Arg	
176	420					425					430					435	
178	tcg	ttc	ttc	gcg	cag	tcg	cag	gtg	ctg	ccg	agg	atc	acc	gac	gcg	ttg	2300
179	Ser	Phe	Phe	Ala	Gln	Ser	Gln	Val	Leu	Pro	Arg	Ile	Thr	Asp	Ala	Leu	
180				440						445				450			
182	gcg	gag	cgc	tgc	tcg	acc	ccg	ctg	ggc	cgg	gcc	acc	tac	ccc	gct	tcc	2348
183	Ala	Glu	Arg	Cys	Ser	Thr	Pro	Leu	Gly	Arg	Ala	Thr	Tyr	Pro	Ala	Ser	
184			455					460					465				
186	ggc	ctc	gcg	gtg	ctg	gac	ggt	tcg	cgg	acg	gac	tgc	gcg	ctg	ggc	agc	2396
187	Gly	Leu	Ala	Val	Leu	Asp	Gly	Ser	Arg	Thr	Asp	Cys	Ala	Leu	Gly	Ser	
188		470					475					480					
190	gac	ccg	gac	gcg	gtg	cgg	ccg	ggg	atc	ttc	ggc	ccg	ggc	cgg	atg	ccg	2444
191	Asp	Pro	Asp	Ala	Val	Arg	Pro	Gly	Ile	Phe	Gly	Pro	Gly	Arg	Met	Pro	
192		485					490					495					
194	gtg	ctg	aag	aac	cag	ccg	tac	gtg	gag	aac	tcc	aac	gac	agc	gcg	tgg	2492
195	Val	Leu	Lys	Asn	Gln	Pro	Tyr	Val	Glu	Asn	Ser	Asn	Asp	Ser	Ala	Trp	
196	500					505					510					515	
198	ctg	acc	aat	gcg	gag	cgg	ccg	ctg	acc	ggg	tac	gag	cgg	gtc	ttc	ggc	2540
199	Leu	Thr	Asn	Ala	Glu	Arg	Pro	Leu	Thr	Gly	Tyr	Glu	Arg	Val	Phe	Gly	
200				520						525					530		
202	acg	atc	gcg	acg	ccc	cgg	tcg	atg	cgg	acg	cgc	ggc	gcg	atc	gag	gac	2588
203	Thr	Ile	Ala	Thr	Pro	Arg	Ser	Met	Arg	Thr	Arg	Gly	Ala	Ile	Glu	Asp	

DATE: 10/19/2004

PATENT APPLICATION: US/10/019.282

TIME: 17:29:38

Input Set : D:\217770US0PCT.txt

Output Set: N:\CRF4\10192004\J019282.raw

204					535				540				545							
206	gtc	gcg	tcg	atg	gcg	gac	cgg	ggc	cgc	ctc	cgg	gtc	ggg	gac	ctt	cag	2636			
207	Val	Ala	Ser	Met	Ala	Asp	Arg	Gly	Arg	Leu	Arg	Val	Gly	Asp	Leu	Gln				
208	550				555				560											
210	cgg	cag	cag	ttc	gcc	aac	cgt	gcg	ccg	gcc	ggg	gat	ctg	gcc	gcc	ttc	2684			
211	Arg	Gln	Gln	Phe	Ala	Asn	Arg	Ala	Pro	Ala	Gly	Asp	Leu	Ala	Ala	Ser				
212	565				570				575											
214	gag	gcc	gcc	aag	tgg	tgt	gcg	gcg	ctg	ccg	ggc	ggc	acg	gcc	gtg	ggc	2732			
215	Glu	Ala	Ala	Lys	Trp	Cys	Ala	Ala	Leu	Pro	Gly	Gly	Thr	Ala	Val	Gly				
216	580				585				590				595							
218	tcc	gac	gga	acg	ccg	gtc	gac	gtg	tcg	gcg	gcc	tgc	cgg	gtg	ctg	cgg	2780			
219	Ser	Asp	Gly	Thr	Pro	Val	Asp	Val	Ser	Ala	Ala	Cys	Arg	Val	Leu	Arg				
220	600				605				610											
222	cgc	tgg	gac	cgg	acc	gtg	gac	agc	gac	agc	cgg	ggc	gcg	ctg	ctc	ttc	2828			
223	Arg	Trp	Asp	Arg	Thr	Val	Asp	Ser	Asp	Ser	Arg	Gly	Ala	Leu	Leu	Phe				
224	615				620				625											
226	gac	cgg	ttc	tgg	cgg	aag	gcg	tcg	tcg	gcg	ccc	gcc	gcc	gag	ctg	tgg	2876			
227	Asp	Arg	Phe	Trp	Arg	Lys	Ala	Ser	Ser	Ala	Pro	Ala	Ala	Glu	Leu	Trp				
228	630				635				640											
230	agg	acg	ccg	ttc	gat	ccg	gcc	gac	ccg	gtg	cgc	act	ccg	cgc	ggc	ctg	2924			
231	Arg	Thr	Pro	Phe	Asp	Pro	Ala	Asp	Pro	Val	Arg	Thr	Pro	Arg	Gly	Leu				
232	645				650				655											
234	aac	acg	gcc	gcg	ccc	gtc	ctg	ggc	agg	gcc	ctg	gcg	gac	gcc	gtg	gcg	2972			
235	Asn	Thr	Ala	Ala	Pro	Val	Leu	Gly	Arg	Ala	Leu	Ala	Asp	Ala	Val	Ala				
236	660				665				670				675							
238	gag	ctg	cgg	gcg	gcg	ggc	atc	gcg	ctg	gac	gcc	ccg	ctg	ggc	gag	cac	3020			
239	Glu	Leu	Arg	Ala	Ala	Gly	Ile	Ala	Leu	Asp	Ala	Pro	Leu	Gly	Glu	His				
240	680				685				690											
242	cag	ttc	gtc	gtg	cgg	aac	ggc	aag	cgg	ctc	ccg	atc	ggc	ggc	ggg	acg	3068			
243	Gln	Phe	Val	Arg	Asn	Gly	Lys	Arg	Leu	Pro	Ile	Gly	Gly	Gly	Thr					
244	695				700				705											
246	gag	tcg	ctg	ggc	atc	tgg	aac	aag	acc	gag	ccg	cag	tgg	aac	gcg	gcg	3116			
247	Glu	Ser	Leu	Gly	Ile	Trp	Asn	Lys	Thr	Glu	Pro	Gln	Trp	Asn	Ala	Ala				
248	710				715				720											
250	ggc	ggc	ggc	tat	acg	gag	gtg	tcg	tcg	ggc	tcc	agc	tac	atc	cag	gcg	3164			
251	Gly	Gly	Gly	Tyr	Thr	Glu	Val	Ser	Ser	Gly	Ser	Ser	Tyr	Ile	Gln	Ala				
252	725				730				735											
254	gtc	ggc	tgg	gac	gac	agc	cgc	tgc	ccg	gtg	gcc	cgg	acg	ctg	ctg	acg	3212			
255	Val	Gly	Trp	Asp	Asp	Ser	Arg	Cys	Pro	Val	Ala	Arg	Thr	Leu	Leu	Thr				
256	740				745				750				755							

RAW SEQUENCE LISTING

DATE: 10/19/2004

PATENT APPLICATION: US/10/019,282

TIME: 17:29:38

Input Set : D:\217770US0PCT.txt

Output Set: N:\CRF4\10192004\J019282.raw

```

270 cgg tag cgcggtgggc ggacgggccc gcccatccgc ggcgagaagg gcgtccgcct 3412
271 Arg
274 cggcggggcgc ccttctcacc gatgtgtcgt gaccgcgctc ccggggggcgt cctcaccgag 3472
276 ccgccgaagg gcccggcggc cgaacccgtg accatgcgtg cgacgcacat ca cgtccgcgtc 3532
278 gctccgccct ccgcccgcgc ccaggccagc tgcgcgtcgc tcagcgggcg gtcgaagcct 3592
280 tccgggaaca gcagcatccg cggctgcggc cacatgttct ccggtccgtg ttcctgacag 3652
282 tccagggcga ggagatgcgg cccgtccccg caggactcgt gccggtaggg gcggtcgtgc 3712
284 gcccggcaga aatagccgaa caccgcacag tggtcgtcgc cgcccggtcg gtggaagccg 3772
286 gggtcgctga cgatcacggc caccggctcc tgccggttga gccgagggat gggccgggga 3832
288 tcacgccaca acagtgcagg agggagcaca cgctcatctt ccccggggcc gagcccacgg 3892
290 gaagggggag cacggcgga cgctcccgt cggcgtgatc gaccggggcg tcccgcgtcg 3952
292 gggcgggccc tcccggacc gtgtctctac agcggggcgt cgaagccctc ccagtacgg 4012
294 tcgcgcagcc gccgtttgta gagcttgccg ttggggtcgc ggggcatggc ggtgatgaag 4072
296 tcgaggctcc ggggtcgttt gtagccggcg agccgctcct cgagtgggc gaggatcgcg 4132
298 gcggcgagcg cgggtgacgg ctggtggcca tcggccgggt cgacgacggc cttgacctcc 4192
300 tcgcccgcgt cgccgtgggg gatgccgaag gcggcgcggt ccgcgacggc ggggtgggtg 4252
302 agcaggaccg actcgatctc ggcggggtag atgttgaccc cgcccgcgat gatcatgtcg 4312
304 atcttgcggt cgcgaggaa gaggtagccg tcctcgtcca gcacgccgag gtaccgacg 4372
306 gtgaagaagt cgccgatgcg gttcgtgcgg gtcttggtct cgtccttggtg gtagctgaag 4432
308 ccgccggtgc tcattctcat gtagacgggtg ccagttcgc ctggcgggag gcggttgccg 4492
310 tcgtcgtcga agacggccag ttcgctgatc ggccaggcct tgccgacggg gccgggcttc 4552
312 ttcagccagt cctcggcggt ggccaacgt ccccgccct cgctggccgc gtagtactcc 4612
314 tcgacgcagc tccccacca gtcgatcatg gcgcgttga cgtggtcggg gcagggggct 4672
316 gcccgtgga tggcgtgccg catggaggag acgtcgtagc gggacctcac ctgctcgggc 4732
318 agcgcgagca gccggtgga ctgggtgggg accatgtggg tgtgggtgca gcggtgggcg 4792
320 tcgacgaggc gcagcatctc ctggggcgac cagccgtcca tcaggaccag cgggtggccg 4852
322 atgtgcaggg cggcgccgc gaattggagt acggcggtgt ggtagagcgg cgagcagacc 4912
324 aggtggacgt tgcgtcgaa cggccggatg ccgaagatgc cgaggaacct gccgaggtag 4972
326 gtctcctcgg ggcgtttgcc gggcaggggg cgccgatgc cgcgggggcg gccggtggtg 5032
328 cccgaggtgt agttcatgac ccagccgagg gtgcggttct caggcgcggt ggcggggtgg 5092
330 ccttcagga gttcggcca gggcgggcag ccggggaccg tgccgacgcc gtagcggtgg 5152
332 gtcgcgggca gttccgctc gtcggcgcg cgcgtcgcgg tggccgcgaa gcgttcgtgg 5212
334 gcgatcagga cgcgggcgcc ggagtcggcg acgatccagg cgatctcggg gccgacgagg 5272
336 tgggtggtga ccggcacgag gtagaagccg gcctgcgagg cggcgaggtg ggcggtgagg 5332
338 agttcgacgc cgttgggcag gacgacggcg aacgcgtcgc ctcgcgcag tccggccgcg 5392
340 cgcaggccgt ggaccatgcg gttgacgtcg gcgtgcaggc ggcccgcgt ccactcctcg 5452
342 ccgtcggggg cgatcaggac ggtgcggtcg ggtcgggtg cggcctgggc ccagaaaccg 5512
344 ttggcggtt ggttcacgtg gcactccttc cggcgatgcg gttcatgcgg gtgacggccc 5572
346 gttcgaagcc gcgggtcagg tcgtcgacga cggcccggac gctgcgttca ctggtcatcc 5632
348 ggccgacgat ctgcccgcg ggcggtgccg gcagctcgcc gacctcgta tcttgatcc 5692
351 <210> SEQ ID NO: 2
352 <211> LENGTH: 804
353 <212> TYPE: PRT
354 <213> ORGANISM: Streptomyces Sp.
356 <400> SEQUENCE: 2
358 Leu Thr Leu Arg Asn Arg Leu Arg Leu Leu Gly Val Ala Gly Leu Ala
359 1 5 10 15
362 Leu Phe Thr Val Ser Ala Ser Leu Pro Pro Ala Thr Ala Ser Gly Thr
363 20 25 30

```

VERIFICATION SUMMARY

DATE: 10/19/2004

PATENT APPLICATION: US/10/019,282

TIME: 17:29:39

Input Set : D:\217770US0PCT.txt

Output Set: N:\CRF4\10192004\J019282.raw